

Figure 1, page 1

EcoR I

- 4102 - 4152 AGGAATTCAAT CCATTTAAAT CATAACAATT AATGGCTTT AGTATATTCA
 - 4052 - 4102 CAGGGTGTGC ATCCATCACA ATCCATTAA GAACAGTTT ATTACTCAA
<HNF-3/Fkh-1
 B2 - 4002 - 4052 AAATAAACCC TGCATTCCTT AGCCATCACCC CCCAACATC CTCCATCCTC
 - 3952 - 4002 CTTCCAAGCC CTGGGCAACC ACCAATCTAC TTTCTGTCTC TATAAATTG
 - 3902 - 3952 CCAATTCTGG ACATTTCATA TAAATGGAAG CAAACAAACAT GTGAGACTT
 - 3852 - 3902 *<NF-Y* GTGACTGGCT GCTTTCACTT AGCATTCTAT TTTAAGGCT CATTATGTTA
 - 3802 - 3852 CAGTACTTAG CAGTACTTCA TTCTTTTTA TTCTCAAATG GTATTCCACT
 - 3752 - 3802 GTGTGGGTAT CCCATATCAT ATTATTAGAG ACAGGTTCTC ACTCTGTAC
 - 3702 - 3752 CCAGGCTGGA GTGCAGTGGC ACAATCATAG CTCACTGTAA CCTCAAAC
 - 3652 - 3702 *<SREBP* CTGGGCTCAA STGATCCTAC TACCTCAGCC TCCAGAGTAG CTAGGACTAC
 - 3602 - 3652 AGGCACACAC AGCCATACCT GGCTAAATT TTTTTTTAAT *<IRF-1*
 - 3552 - 3602 TGTTTCATT TTCTTTCTT TTTGTTGTTG TTGTTTGAG ATAGGGTCTC
 - 3502 - 3552 ACTTTGTTAC CCAGGCTGGA GGGCAGTGGC ATGGTGACAG CTGAGCAGCC
 - 3452 - 3502 *<SREBP* TTGACTTCCT GGGCTCAAGT GATCCTCCTG CCTCAGCCTC CCAAGTAGCT
 - 3402 - 3452 CGGACTACAA ACACGTGTCA CCATGCCTGG CTGATATT TTTCTTGAA
 - 3352 - 3402 ACAGGGTATC ACTCTGTGCA CCAGGCTGGA GTACAGTGGC GTATAATAG
c1
 Pst I

- 3302 - 3352 CTCACTGCAG CCTCCCCCTCC TGGGCTCAAG CAATCCGCTG GCCTCAGCAT
 - 3252 - 3302 CCTGAGTAGC TGGGACTACA GGCTGTGCC ACCAGGCCA GCTAAGTTT
 - 3202 - 3252 AAAAATGAT TTTGGTATA GAGGAGGTCT TGCTATGTTG CTCAGGCTGT
 - 3152 - 3202 ATTTTATTG TTGAGACAAG GTCTCACTAT GTGCCATGA *<AP-1* TCCCCCACC
 - 3102 - 3152 TCCACTTCCC AAAGTGCTCA TCTTATCTGT TCATTAGTCA GTTGACAGAC
 - 3052 - 3102 ATTTAGGTTG TTTCCACTT TTGACCAATTAA TGAATAATAC TCCAGTGAAT
 - 3002 - 3052 ATTCATGTAT ACATTGTGT GGGCATAATGT TTTCATTCT GTGGGTTA
 - 2952 - 3002 TATCTAGGAG TGGATTGCT GGATCCGGG TAATATTTG ACAGGCAGAG
 - 2902 - 2952 *C/EBP-β* TTCAGGGGAA GAAAAACTTG GGAAAATGAA GCATGTTAG AAATCAGCAA
 - 2852 - 2902 GAGTGCAGGG GTTTTCGGA GTTTTATTG ATATTCTGTT GACAAATGTG
 - 2752 - 2802 CAGTTGATG AAGATACAAG TTATACTAAG TGAGAAGTGA GAATTAAGGC
 - 2702 - 2752 TGGAATAGGG CGTCAGAGT AAAATCATGA AGCACTTGA ATACCAAAAT
 - 2702 - 2752 *NF-1* *<HNF3-β* TAAGGAGCTT GGCTGTAAAC AAAATAATAA AAAATCACAA TTTTTTTT
 - 2652 - 2702 TTTTTGAGA AAGAGTCTTG CTCTTCACC CTGGCTGGAG GGCAGTGGTG
 - 2602 - 2652 *<SREBP* TGATCTCAGC TCACTGCAAC TTTGCCCTCC CGGGTTCAAG CAATTCTCCT

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-2552 -2602 GCTTCAGCCT CCCAAGTAGC TGGGACTACA GGCACTTCCC ACCATGCCCA <NF-kB
 -2502 -2552 GCTGATTTT GTATTTTAG TAGAGATGGG ATTC CACTTT GTTGGCCAAG NF-kB <IRF-1
 -2452 -2502 CTGGTCTCAA ACTTTTGCT GTCATAATTG TTGTAACAT ATGTCCTTT
 -2402 -2452 GCTGAGGTAG GGCCCC CAGA CCAAAAAAAA TAA ATCTTAG AATCCAAATC AP-2s <HNF3-β
 -2352 -2402 AGTGTGTTGG TTTGACCACT GTCAC TGAG AACCACAGTG TGACCAAGGGC
 B2
 C2
 Taq I
 ---- IRF-2s
 -2302 -2352 CTCAGGAGTA GAGGTGATCT CTGCTCGAAA GAGAA ATAGA ATGAAAATAT
 -2252 -2302 TCTCCGGGCC AGGCGTGGTG GCTCATGCCT GTAATCCCAG CACTITGGGA
 T3R
 -2202 -2252 GCCCAAGGCA TGTGGATCAC CTGAGGTCAG GAGTTCAAAA CCAGCCTGGC SREBP > RAR-α1
 -2152 -2202 CAACATGGTG AAACCCGTC TCTACTAAAA ATACAAAAAA TTAGCTAAGT
 -2102 -2152 GTGGTGGCCG ATGCCTGTAA TCCCAGCTAC TTGGGAGGGT GAGGCAGGAG
 -2052 AATTCTTGA ACCCGGGAGG CAGAGGTTGC AGTGAAGCGA GATCAC ACCA <Pax-6 SREBP>
 -2002 CTGCACTCCA GCCTGGGG GAGAGCGAGA CTTCCCTCTCA AAAAAA CAAA <HNF-3/Fkh-2
 -1952 AAACAAAAGA ATTAAGCAA TTAGACATTG CAGAGAGAAC CTGAAGGGGG C/EBP-β <CHOP>
 RAR-α1>
 -1902 TCAGACCACG TACAGATTC TGTGCCACAT GCC AA GTACT TCTGAGGC AT <NF-1 Pax-6>
 -1852 GACTGGATGA GCTGTCCACA TCTGAAATCA TCCAGTCTTG TTCAGAACCTT
 -1802 TCACACCGGA CAGGGAGCCA GGACTGGAAT GCAGTCTCCT GGT CA CTGGC RAR-α1
 NF-1 > <ER> <NF-1>
 -1752 CAGAGAGTTG GCCT TGACCC TGAGACCAGT GGCC AA CAAA GGAGCTGCTT
 <Rel>
 -1702 AGTCTACCTC CCAGGAA ATC CCAGGTGCTT GTCTCCTGG GAAGTGA ATC AP-1
 <NF-1>
 -1652 ATGGCG CAG CACTCCGTAT TTTCTCCTCT TCCCAGGGGA AGGATCCTAG
 -1602 GGCAGTATTG GGGAAAGACA TGGGCATGGGA AGGAC ACCCG GTGAATGCAT <GR>
 C3
 Sac I

 -1552 AGCCTGCCTG GTTCTGAGCT CTCATGGTAA GGCTCCTACA GACACGGAAA
 -1502 AGATGGGGC ACAGGGACAG ATCAGTAGGG TCAGAGCATC TCAGGGACCG
 -1452 AGGGCAATAT GGTCTGAGC AGGGATTAAG AGCTTGGCT CTCATATGGT
 <CREB> <ER>
 -1402 GTTTCTGGC TCAACTGCCA GCTCCGT CAC TTACTGGTG CTGTGAC CAT
 -1352 GGGCAAGTTA TTCCATCTCT CCATATCTCT TTCCACTT TTAAAATGGA